

PCT/10 W

CRF Errors Corrected by the STIC Systems Branch

Serial Number: 10/069,626

CRF Processing Date: 11/6/2002  
Edited by: AN  
Verified by: AN (STIC staff)

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☐ Other: \_\_\_\_\_

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

75



PCT10

## RAW SEQUENCE LISTING

DATE: 11/06/2002

PATENT APPLICATION: US/10/069,626

TIME: 18:25:51

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\11062002\J069626.raw

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6   Lymphocyte Activation Antigen B-7 Family and
7   Polypeptides Encoded Thereby
9 <130> FILE REFERENCE: 15966-562 NATL
11 <140> CURRENT APPLICATION NUMBER: 10/069,626
C--> 12 <141> CURRENT FILING DATE: 2002-07-25
14 <150> PRIOR APPLICATION NUMBER: PCT/US00/24220
15 <151> PRIOR FILING DATE: 2000-08-31
17 <150> PRIOR APPLICATION NUMBER: 60/152383
18 <151> PRIOR FILING DATE: 1999-09-03
20 <150> PRIOR APPLICATION NUMBER: 60/172909
21 <151> PRIOR FILING DATE: 1999-12-21
23 <150> PRIOR APPLICATION NUMBER: 60/183578
24 <151> PRIOR FILING DATE: 2000-02-18
26 <150> PRIOR APPLICATION NUMBER: 09/651200
27 <151> PRIOR FILING DATE: 2000-08-30
29 <160> NUMBER OF SEQ ID NOS: 25
31 <170> SOFTWARE: PatentIn Ver. 2.0
33 <210> SEQ ID NO: 1
34 <211> LENGTH: 2691
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46                                     Met Ala
47                                     1
49 aac gag cag ggc ttg ttt gat gtg cac agc atc ctg cgg gtg gtg ctg 164
50 Asn Glu Gln Gly Leu Phe Asp Val His Ser Ile Leu Arg Val Val Leu
51      5              10              15
53 ggt gca aat ggc acc tac agc tgc ctg gtg cgc aac ccc gtg ctg cag 212
54 Gly Ala Asn Gly Thr Tyr Ser Cys Leu Val Arg Asn Pro Val Leu Gln
55      20              25              30
57 cag gat gcg cac agc tct gtc acc atc aca ccc cag aga agc ccc aca 260
58 Gln Asp Ala His Ser Ser Val Thr Ile Thr Pro Gln Arg Ser Pro Thr
59      35              40              45              50
61 gga gcc gtg gag gtc cag gtc cct gag gac ccg gtg gtg gcc cta gtg 308
62 Gly Ala Val Glu Val Gln Val Pro Glu Asp Pro Val Val Ala Leu Val
63      55              60              65

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69 agc ctg aca cag ctc aac ctc atc tgg cag ctg aca gac acc aaa cag 404
70 Ser Leu Thr Gln Leu Asn Leu Ile Trp Gln Leu Thr Asp Thr Lys Gln
71          85          90          95
73 ctg gtg cac agt ttc acc gaa ggc cgg gac cag ggc agc gcc tat gcc 452
74 Leu Val His Ser Phe Thr Glu Gly Arg Asp Gln Gly Ser Ala Tyr Ala
75          100          105          110
77 aac cgc acg gcc ctc ttc ccg gac ctg ctg gca caa ggc aat gca tcc 500
78 Asn Arg Thr Ala Leu Phe Pro Asp Leu Leu Ala Gln Gly Asn Ala Ser
79 115          120          125          130
81 ctg agg ctg cag cgc gtg cgt gtg gcg gac gag ggc agc ttc acc tgc 548
82 Leu Arg Leu Gln Arg Val Arg Val Ala Asp Glu Gly Ser Phe Thr Cys
83          135          140          145
85 ttc gtg agc atc cgg gat ttc ggc agc gct gcc gtc agc ctg cag gtg 596
86 Phe Val Ser Ile Arg Asp Phe Gly Ser Ala Ala Val Ser Leu Gln Val
87          150          155          160
89 gcc gct ccc tac tgc aag ccc agc atg acc ctg gag ccc aac aag gac 644
90 Ala Ala Pro Tyr Ser Lys Pro Ser Met Thr Leu Glu Pro Asn Lys Asp
91          165          170          175
93 ctg cgg cca ggg gac acg gtg acc atc acg tgc tcc agc tac cgg ggc 692
94 Leu Arg Pro Gly Asp Thr Val Thr Ile Thr Cys Ser Ser Tyr Arg Gly
95          180          185          190
97 tac cct gag gct gag gtg ttc tgg cag gat ggg cag ggt gtg ccc ctg 740
98 Tyr Pro Glu Ala Glu Val Phe Trp Gln Asp Gly Gln Gly Val Pro Leu
99 195          200          205          210
101 act ggc aac gtg acc acg tgc cag atg gcc aac gag cag ggc ttg ttt 788
102 Thr Gly Asn Val Thr Thr Ser Gln Met Ala Asn Glu Gln Gly Leu Phe
103          215          220          225
105 gat gtg cac agc gtc ctg cgg gtg gtg ctg ggt gcg aat ggc acc tac 836
106 Asp Val His Ser Val Leu Arg Val Val Leu Gly Ala Asn Gly Thr Tyr
107          230          235          240
109 agc tgc ctg gtg cgc aac ccc gtg ctg cag cag gat gcg cac ggc tct 884
110 Ser Cys Leu Val Arg Asn Pro Val Leu Gln Gln Asp Ala His Gly Ser
111          245          250          255
113 gtc acc atc aca ggg cag cct atg aca ttc ccc cca gag gcc ctg tgg 932
114 Val Thr Ile Thr Gly Gln Pro Met Thr Phe Pro Pro Glu Ala Leu Trp
115          260          265          270
117 gtg acc gtg ggg ctc tct gtc tgt ctc att gca ctg ctg gtg gcc ctg 980
118 Val Thr Val Gly Leu Ser Val Cys Leu Ile Ala Leu Leu Val Ala Leu
119 275          280          285          290
121 gct ttc gtg tgc tgg aga aag atc aaa cag agc tgt gag gag gag aat 1028
122 Ala Phe Val Cys Trp Arg Lys Ile Lys Gln Ser Cys Glu Glu Glu Asn
123          295          300          305
125 gca gga gcc gag gac cag gat ggg gag gga gaa ggc tcc aag aca gcc 1076
126 Ala Gly Ala Glu Asp Gln Asp Gly Glu Gly Glu Gly Ser Lys Thr Ala
127          310          315          320
129 ctg cag cct ctg aaa cac tct gac agc aaa gaa gat gat gga caa gaa 1124

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130 Leu Gln Pro Leu Lys His Ser Asp Ser Lys Glu Asp Asp Gly Gln Glu
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133 ata gcc tgaccatgag gaccagggag ctgctacccc tccctacagc tcctaccctc 1180
134 Ile Ala
135          340
137 tggctgcaat ggggctgcac tgtgagccct gcccccaaca gatgcatcct gctctgacag 1240
139 gtgggctcct tctccaaagg atgcgataca cagaccactg tgcagcctta tttctccaat 1300
141 ggacatgatt cccaagtcac cctgctgcct tttttcttat agacacaatg aacagaccac 1360
143 ccacaacctt agttctctaa gtcacccctg ctgctgcctt atttcacagt acatacatTT 1420
145 cttagggaca cagtacactg accacatcac caccctcttc ttccagtgtc gcgtggacca 1480
147 tctggctgcc ttttttctcc aaaagatgca atattcagac tgactgaccc cctgccttat 1540
149 ttcaccaaag acacgatgca tagtcacccc ggccttggtt ctccaatggc cgtgatacac 1600
151 tagtgatcat gttcagccct gcttcacact gcatagaatc ttttcttctc agacagggac 1660
153 agtgcgccct caacatctcc tggagtctag aagctgtttc ctttcccttc cttcctcctc 1720
155 ttgctctagc cttaatactg gccttttccc tccctgcccc aagtgaagac agggcactct 1780
157 gcgcccacca catgcacagc tgtgcatgga gacctgcagg tgcacgtgct ggaacacgtg 1840
159 tggttccccc ctggcccagc ctctcttgca gtgcccctct cccctgcccc tcctccccac 1900
161 ggaagcatgt gctgggtcac ctggttctcc aggggtctgt gatggggccc ctgggggtca 1960
163 gcttctgtcc ctctgccttc tcacctttt gtctctttct tttcatgtat ccattcagtt 2020
165 gatgtttatt gagcaactac agatgtcagc actgtgttag gtgctggggg ccttgcgtgg 2080
167 gaagataaag ttctccctc aaggactccc catccagctg ggagacagac aactaactac 2140
169 actgcaccct gcggtttgca gggggctcct gcctggctcc ctgctccaca cctcctctgt 2200
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187 ctttgtaaaa a 2691
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191 <211> LENGTH: 340
192 <212> TYPE: PRT.
193 <213> ORGANISM: Homo sapiens
195 <400> SEQUENCE: 2
196 Met Ala Asn Glu Gln Gly Leu Phe Asp Val His Ser Ile Leu Arg Val
197 1 5 10 15
199 Val Leu Gly Ala Asn Gly Thr Tyr Ser Cys Leu Val Arg Asn Pro Val
200 20 25 30
202 Leu Gln Gln Asp Ala His Ser Ser Val Thr Ile Thr Pro Gln Arg Ser
203 35 40 45
205 Pro Thr Gly Ala Val Glu Val Gln Val Pro Glu Asp Pro Val Val Ala
206 50 55 60
208 Leu Val Gly Thr Asp Ala Thr Leu His Cys Ser Phe Ser Pro Glu Pro
209 65 70 75 80
211 Gly Phe Ser Leu Thr Gln Leu Asn Leu Ile Trp Gln Leu Thr Asp Thr
212 85 90 95
214 Lys Gln Leu Val His Ser Phe Thr Glu Gly Arg Asp Gln Gly Ser Ala

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215          100          105          110
217 Tyr Ala Asn Arg Thr Ala Leu Phe Pro Asp Leu Leu Ala Gln Gly Asn
218          115          120          125
220 Ala Ser Leu Arg Leu Gln Arg Val Arg Val Ala Asp Glu Gly Ser Phe
221          130          135          140
223 Thr Cys Phe Val Ser Ile Arg Asp Phe Gly Ser Ala Ala Val Ser Leu
224 145          150          155          160
226 Gln Val Ala Ala Pro Tyr Ser Lys Pro Ser Met Thr Leu Glu Pro Asn
227          165          170          175
229 Lys Asp Leu Arg Pro Gly Asp Thr Val Thr Ile Thr Cys Ser Ser Tyr
230          180          185          190
232 Arg Gly Tyr Pro Glu Ala Glu Val Phe Trp Gln Asp Gly Gln Gly Val
233          195          200          205
235 Pro Leu Thr Gly Asn Val Thr Thr Ser Gln Met Ala Asn Glu Gln Gly
236          210          215          220
238 Leu Phe Asp Val His Ser Val Leu Arg Val Val Leu Gly Ala Asn Gly
239 225          230          235          240
241 Thr Tyr Ser Cys Leu Val Arg Asn Pro Val Leu Gln Gln Asp Ala His
242          245          250          255
244 Gly Ser Val Thr Ile Thr Gly Gln Pro Met Thr Phe Pro Pro Glu Ala
245          260          265          270
247 Leu Trp Val Thr Val Gly Leu Ser Val Cys Leu Ile Ala Leu Leu Val
248          275          280          285
250 Ala Leu Ala Phe Val Cys Trp Arg Lys Ile Lys Gln Ser Cys Glu Glu
251          290          295          300
253 Glu Asn Ala Gly Ala Glu Asp Gln Asp Gly Glu Gly Glu Gly Ser Lys
254 305          310          315          320
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268 <220> FEATURE:
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270 <222> LOCATION: (2)..(1324)
272 <400> SEQUENCE: 3
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274 Pro Leu Pro Gly Pro Ala Gly Thr Gly Gln Arg Ile Pro Glu Ala Ala
275 1 5 10 15
277 gcg cgt gcg tgt agc gga cga ggg cag ctt cac ctg ctt cgt gag cat 97
278 Ala Arg Ala Cys Ser Gly Arg Gly Gln Leu His Leu Leu Arg Glu His
279 20 25 30
281 ccg gga ttt cgg cag cgc tgc cgt cag cct gca ggt ggc cgc tcc cta 145
282 Pro Gly Phe Arg Gln Arg Cys Arg Gln Pro Ala Gly Gly Arg Ser Leu
283 35 40 45
285 ctc gaa gcc cag cat gac cct gga gcc caa caa gga cct gcg gcc agg 193

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## RAW SEQUENCE LISTING

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TIME: 18:25:51

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\11062002\J069626.raw

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286 Leu Glu Ala Gln His Asp Pro Gly Ala Gln Gln Gly Pro Ala Ala Arg
287      50      55      60
289 gga cac ggt gtg acc atc acg tgc tcc agc tac cag ggc tac cct gag 241
290 Gly His Gly Val Thr Ile Thr Cys Ser Ser Tyr Gln Gly Tyr Pro Glu
291 65      70      75      80
293 gct gag gtg ttc tgg cag gat ggg cag ggt gtg ccc ctg act ggc aac 289
294 Ala Glu Val Phe Trp Gln Asp Gly Gln Gly Val Pro Leu Thr Gly Asn
295      85      90      95
297 gtg acc acg tcg cag atg gcc aac gag cag ggc ttg ttt gat gtg cac 337
298 Val Thr Thr Ser Gln Met Ala Asn Glu Gln Gly Leu Phe Asp Val His
299      100      105      110
301 agc atc ctg cgg gtg gtg ctg ggt gca aat ggc acc tac agc tgc ctg 385
302 Ser Ile Leu Arg Val Val Leu Gly Ala Asn Gly Thr Tyr Ser Cys Leu
303      115      120      125
305 gtg cgc aac ccc gtg ctg cag cag gat gcg cac agc tct gtc acc atc 433
306 Val Arg Asn Pro Val Leu Gln Gln Asp Ala His Ser Ser Val Thr Ile
307      130      135      140
309 aca ccc cag aga agc ccc aca gga gcc gtg gag gtc cag gtc cct gag 481
310 Thr Pro Gln Arg Ser Pro Thr Gly Ala Val Glu Val Gln Val Pro Glu
311 145      150      155      160
313 gac ccg gtg gtg gcc cta gtg ggc acc gat gcc acc ctg cac tgc tcc 529
314 Asp Pro Val Val Ala Leu Val Gly Thr Asp Ala Thr Leu His Cys Ser
315      165      170      175
317 ttc tcc ccc gag cct ggc ttc agc ctg aca cag ctc aac ctc atc tgg 577
318 Phe Ser Pro Glu Pro Gly Phe Ser Leu Thr Gln Leu Asn Leu Ile Trp
319      180      185      190
321 cag ctg aca gac acc aaa cag ctg gtg cac agt ttc acc gaa ggc cgg 625
322 Gln Leu Thr Asp Thr Lys Gln Leu Val His Ser Phe Thr Glu Gly Arg
323      195      200      205
325 gac cag ggc agc gcc tat gcc aac cgc acg gcc ctc ttc ccg gac ctg 673
326 Asp Gln Gly Ser Ala Tyr Ala Asn Arg Thr Ala Leu Phe Pro Asp Leu
327      210      215      220
329 ctg gca caa ggc aat gca tcc ctg agg ctg cag cgc gtg cgt gtg gcg 721
330 Leu Ala Gln Gly Asn Ala Ser Leu Arg Leu Gln Arg Val Arg Val Ala
331 225      230      235      240
333 gac gag ggc agc ttc acc tgc ttc gtg agc atc cgg gat ttc ggc agc 769
334 Asp Glu Gly Ser Phe Thr Cys Phe Val Ser Ile Arg Asp Phe Gly Ser
335      245      250      255
337 gct gcc gtc agc ctg cag gtg gcc gct ccc tac tcg aag ccc agc atg 817
338 Ala Ala Val Ser Leu Gln Val Ala Ala Pro Tyr Ser Lys Pro Ser Met
339      260      265      270
341 acc ctg gag ccc aac aag gac ctg cgg cca ggg gac acg gtg acc atc 865
342 Thr Leu Glu Pro Asn Lys Asp Leu Arg Pro Gly Asp Thr Val Thr Ile
343      275      280      285
345 acg tgc tcc agc tac cgg ggc tac cct gag gct gag gtg ttc tgg cag 913
346 Thr Cys Ser Ser Tyr Arg Gly Tyr Pro Glu Ala Glu Val Phe Trp Gln
347      290      295      300
349 gat ggg cag ggt gtg ccc ctg act ggc aac gtg acc acg tcg cag atg 961
350 Asp Gly Gln Gly Val Pro Leu Thr Gly Asn Val Thr Thr Ser Gln Met

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RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/069,626

DATE: 11/06/2002  
TIME: 18:25:52

Input Set : A:\PTO.AMC.txt  
Output Set: N:\CRF4\11062002\J069626.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

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RAW SEQUENCE LISTING ERROR SUMMARY  
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Seq#:5; Line(s) 695,696  
Seq#:6; Line(s) 697,698,699,700,701,702,703,704,705,706,707,708,709,710,711  
Seq#:6; Line(s) 712,713,714,715,716,717,718,719,720,721,722,723,724,725,726  
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Seq#:6; Line(s) 802,803,804,805  
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Seq#:8; Line(s) 907  
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Seq#:10; Line(s) 1014,1015,1016,1017,1018,1019,1020,1021  
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Seq#:11; Line(s) 1066,1067,1068,1069,1070,1071,1072,1073,1074,1075,1076  
Seq#:11; Line(s) 1077,1078,1079,1080,1081,1082  
Seq#:12; Line(s) 1083,1084,1085,1086,1087,1088,1089,1090,1091,1092,1093  
Seq#:12; Line(s) 1094,1095,1096,1097,1098,1099,1100,1101,1102,1103,1104  
Seq#:12; Line(s) 1105,1106,1107,1108,1109  
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Seq#:14; Line(s) 1218,1219,1220,1221,1222,1223,1224,1225,1226,1227,1228



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Seq#:14; Line(s) 1229,1230,1231,1232,1233,1234,1235,1236,1237  
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Seq#:15; Line(s) 1249,1250,1251,1252,1253,1254,1255,1256,1257,1258,1259  
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Seq#:15; Line(s) 1271,1272,1273,1274,1275,1276,1277,1278,1279,1280,1281  
Seq#:15; Line(s) 1282,1283,1284,1285,1286,1287,1288,1289,1290,1291,1292  
Seq#:15; Line(s) 1293,1294,1295,1296,1297,1298,1299,1300,1301  
Seq#:16; Line(s) 1302,1303,1304,1305,1306,1307,1308,1309,1310,1311,1312  
Seq#:16; Line(s) 1313,1314,1315,1316,1317,1318,1319,1320,1321,1322,1323  
Seq#:16; Line(s) 1324,1325,1326,1327,1328,1329,1330,1331,1332,1333,1334  
Seq#:16; Line(s) 1335,1336,1337,1338,1339,1340,1341,1342,1343,1344,1345  
Seq#:16; Line(s) 1346,1347,1348,1349,1350,1351,1352,1353,1354,1355,1356  
Seq#:16; Line(s) 1357,1358,1359,1360,1361,1362,1363,1364,1365  
Seq#:17; Line(s) 1366,1367,1368,1369,1370,1371,1372,1373,1374,1375,1376  
Seq#:17; Line(s) 1377,1378,1379,1380,1381,1382,1383,1384,1385,1386,1387  
Seq#:17; Line(s) 1388,1389,1390,1391,1392,1393,1394,1395,1396,1397,1398  
Seq#:17; Line(s) 1399,1400,1401,1402,1403,1404,1405,1406,1407,1408,1409  
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Seq#:18; Line(s) 1433,1434,1435,1436,1437,1438,1439,1440,1441,1442,1443  
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Seq#:18; Line(s) 1499,1500,1501,1502  
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Seq#:21; Line(s) 1709,1710,1711,1712  
Seq#:22; Line(s) 1713,1714,1715,1716,1717,1718,1719,1720,1721,1722,1723

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Seq#:22; Line(s) 1779  
Seq#:23; Line(s) 1780,1781,1782,1783,1784,1785,1786,1787,1788,1789,1790  
Seq#:23; Line(s) 1791,1792,1793,1794,1795,1796,1797,1798,1799,1800,1801  
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Seq#:23; Line(s) 1835,1836,1837,1838,1839,1840,1841,1842,1843  
Seq#:24; Line(s) 1844,1845,1846,1847,1848,1849,1850,1851,1852,1853,1854  
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Seq#:24; Line(s) 1877,1878,1879,1880,1881,1882,1883,1884,1885,1886,1887  
Seq#:24; Line(s) 1888,1889,1890,1891,1892,1893,1894,1895,1896,1897,1898  
Seq#:24; Line(s) 1899,1900,1901,1902,1903,1904,1905,1906,1907,1908,1909  
Seq#:24; Line(s) 1910,1911,1912,1913,1914,1915,1916,1917,1918,1919,1920  
Seq#:24; Line(s) 1921,1922,1923,1924,1925,1926,1927,1928,1929,1930,1931  
Seq#:24; Line(s) 1932,1933,1934,1935,1936,1937,1938,1939,1940,1941,1942  
Seq#:24; Line(s) 1943,1944,1945,1946,1947,1948,1949,1950,1951,1952,1953  
Seq#:24; Line(s) 1954,1955,1956  
Seq#:25; Line(s) 1957,1958,1959,1960,1961,1962,1963,1964,1965,1966,1967  
Seq#:25; Line(s) 1968,1969,1970,1971,1972,1973,1974,1975,1976,1977,1978  
Seq#:25; Line(s) 1979,1980,1981,1982,1983,1984,1985,1986,1987,1988,1989  
Seq#:25; Line(s) 1990,1991,1992,1993,1994,1995,1996,1997,1998,1999,2000  
Seq#:25; Line(s) 2001,2002,2003,2004,2005,2006,2007,2008,2009,2010,2011  
Seq#:25; Line(s) 2012,2013,2014,2015,2016,2017,2018,2019,2020,2021,2022  
Seq#:25; Line(s) 2023,2024,2025,2026,2027,2028,2029

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L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date